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<110> Gurney, Mark E.
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Pauley, Adele M.
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Lys Arg Arg Lys Val Ile Lys Pro Gly Phe Ile His Ser Pro Trp Lys

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Tyr Ser Leu Gln Phe Asp Gly Ile His Val Val Ser Gly Ser Leu Asp 465 470 475 480

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GIN	Cys	ьeu		THE	ьеu	GIII	GIŞ		ASII	гÃг	nıs	GIII		Ala	Val
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Gly	Lys	Cys	Leu 260		Thr	Leu	Val	Gly 265	His	Thr	Gly	Gly	Val 270	Trp	Ser
Ser	Gln	Met 275	_	Asp	Asn	Ile	Ile 280	Ile	Ser	Gly	Ser	Thr 285	Asp	Arg	Thr
Leu	Lys 290	Val	Trp	Asn	Ala	Glu 295		Gly	Glu	Cys	Ile		Thr	Leu	Tyr

His Met Met Gln Val Ile Glu Pro Gln Phe Gln Arg Asp Phe Ile Ser

(Gly	His	Thr	Ser	Thr	Val	Arg	Cys	Met	His	Leu	His	Glu	Lys	Arg	Val
	305					310					315					320
,	Val	Ser	Gly	Ser	`Arg	Asp	Ala	Thr	Leu	Arg	Val	Trp	Asp	Ile	Glu	Thr
			_		325	•				330					335	
					323					330					333	
			_	_											_	_
•	GТУ	GIn	Суѕ		His	Val	Leu	Met		His	Va⊥	Ala	Ala		Arg	Cys
				340					345					350		
	Val	Gln	Tyr	Asp	Gly	Arg	Arg	Val	Val	Ser	Gly	Ala	Tyr	Asp	Phe	Met
			355					360					365			
	Val	Lys	Val	Trp	Asp	Pro	Glu	Thr	Glu	Thr	Cys	Leu	His	Thr	Leu	Gln
		370					375					380				
	Glv	Hic	Thr	Δen	Δra	Val	Tur	Ser	T.e.11	Gln	Dhe	Asn	Glv	Tle	Hic	Va1
	385	1120	****	11011	111.9	390	-1-	501	Lou		395	1100	011			
	303					330					323					400
							_								_	
	Val	Ser	Gly	Ser	Leu	Asp	Thr	Ser	Ile	Arg	Val	Trp	Asp	Val		Thr
					405					410					415	
	Gly	Asn	Cys	Ile	His	Thr	Leu	Thr	Gly	His	Gln	Ser	Leu	Thr	Ser	Gly
				420					425					430		
	Met	Glu	Leu	Lys	Asp	Asn	Ile	Leu	Val	Ser	Gly	Asn	Ala	Asp	Ser	Thr
			435					440					445			
	Val	Lve	Tle	Trn	Δen	Tle	Lvs	Thr	Glv	Gln	Cve	Len	Gln	Thr	T.eu	Gln
	vai	_	110	. p	пор	110		1	CIJ	0111	CYD		0111	-111	Lea	0111
		450					455					460				
	Gly	Pro	Asn	Lys	His	Gln	Ser	Ala	Val	Thr	Суѕ	Leu	Gln	Phe	Asn	Lys
	465					470					475					480
	Asn	Phe	Val	Ile	Thr	ser	Ser	Asp	Asp	Gly	Thr	Val	Lys	Leu	Trp	Asp
					485					490					495	

Leu Lys Thr Gly Glu Phe Ile Arg Asn Leu Val Thr Leu Glu Ser Gly Gly Ser Gly Gly Val Val Trp Arg Ile Arg Ala Ser Asn Thr Lys Leu Val Cys Ala Val Gly Ser Arg Asn Gly Thr Glu Glu Thr Lys Leu Leu Val Leu Asp Phe Asp Val Asp Met Lys <210> 6 <211> 545 <212> PRT <213> Homo sapiens <400> 6 Met Ile Phe Tyr Lys Met Lys Arg Lys Leu Asp His Gly Ser Glu Val Arg Ser Phe Ser Leu Gly Lys Lys Pro Cys Lys Val Ser Glu Tyr Thr Ser Thr Thr Gly Leu Val Pro Cys Ser Ala Thr Pro Thr Thr Phe Gly Asp Leu Arg Ala Ala Asn Gly Gln Gly Gln Gln Arg Arg Ile Thr Ser Val Gln Pro Pro Thr Gly Leu Gln Glu Trp Leu Lys Met Phe Gln Ser Trp Ser Gly Pro Glu Lys Leu Leu Ala Leu Asp Glu Leu Ile Asp

Ser	Cys	Glu	Pro	Thr	Gln	Val	Lys	His	Met	Met	Gln	Val	Ile	Glu	Pro
			100					105					110		
Gln	Phe	Gln	Arg	Asp	Phe	Ile	Ser	Leu	Leu	Pro	Lys	Glu	Leu	Ala	Leu
		115					120					125			
Tyr	Val	Leu	Ser	Phe	Leu	Glu	Pro	Lys	Asp	Leu	Leu	Gln	Ala	Ala	Gln
	130					135					140				
Thr	Cys	Arg	Tyr	Trp	Arg	Ile	Leu	Ala	Glu	Asp	Asn	Leu	Leu	Trp	Arg
145					150					155					160
Glu	Lys	Cys	Lys	Glu	Glu	Gly	Ile	Asp	Glu	Pro	Leu	His	Ile	Lys	Arg
				165					170					175	
Arg	Lys	Va1	Ile	Lys	Pro	Gly	Phe	Ile	His	Ser	Pro	Trp	Lys	Ser	Ala
			180					185					190		
Tyr	Ile	Arg	Gln	His	Arg	Ile	Asp	Thr	Asn	Trp	Arg	Arg	Gly	Glu	Leu
		195					200					205			
Lys	Ser	Pro	Lys	Val	Leu	Lys	Gly	His	Asp	Asp	His	Val	Ile	Thr	Cys
	210					215					220				
Leu	Gln	Phe	Cys	Gly	Asn	Arg	Ile	Val	Ser	Gly	Ser	Asp	Asp	Asn	Thr
225					230					235					240
Leu	Lys	Val	Trp	Ser	Ala	Va1	Thr	Gly	Lys	Cys	Leu	Arg	Thr	Leu	Val
				245					250					255	
Gly	His	Thr	Gly	Gly	Va1	Trp	Ser	Ser	Gln	Met	Arg	Asp	Asn	Ile	Ile
			260					265					270		
Ile	Ser	Gly	Ser	Thr	Asp	Arg	Thr	Leu	Lys	Val	Trp	Asn	Ala	Glu	Thr
		275					280		•		_	295			

Gly	Glu	Cys	Ile	His	Thr	Leu	Tyr	Gly	His	Thr	Ser	Thr	Val	Arg	Cys
	290					295					300				
Met	His	Leu	His	Glu	Lys	Arg	Va1	Val	Ser	Gly	Ser	Arg	Asp	Ala	Thr
305					310					315					320
T 011	2~~	17-1	m.co	7.55	Ile	C1.,	ωb ν.	C1	C1 m	Cvc	Lou	ui c	17 - 1	T 011	Mot
Leu	AIG	vai	iip		116	Giu	1111	GIY		Суз	пеп	1113	vai		mec
				325					330					335	
Gly	His	Val	Ala	Ala	Val	Arg	Cys	Va1	Gln	Tyr	Asp	Gly	Arg	Arg	Val
			340					345					350		
Val	Ser	Gly	Ala	Tyr	Asp	Phe	Met	Va1	Lys	Val	Trp	Asp	Pro	Glu	Thr
		355					360					365			
Glu	Thr	Cve	Leu	His	Thr	Len	Gln	Glv	Hic	Thr	Asn	Ara	Va 1	ጥህዮ	Ser
Olu	370	Cyb	Lea	1115		375	0111	Cly		****		9	V 41	-7-	501
	3/0					3/3					380				
Leu	Gln	Phe	Asp	Gly	Ile	His	Val	Val	Ser	Gly	Ser	Leu	Asp	Thr	Ser
385					390					395					400
Ile	Arg	Val	Trp	Asp	Val	Glu	Thr	Gly	Asn	Cys	Ile	His	Thr	Leu	Thr
				405					410					415	
Gly	His	Gln	Ser	Leu	Thr	Ser	Gly	Met	Glu	Leu	Lys	Asp	Asn	Ile	Leu
			420					425					430		
17-1	Com	G1	3.00	71-	3 an	Com	⊞b ∞	17-1	T	T10	m~~	2 000	T10	T	mb~
vai	ser	_	ASII	АІА	Asp	ser		Val	гур	116	пр		me	гуу	THE
		435					440					445			
Gly	Gln	Cys	Leu	Gln	Thr	Leu	Gln	Gly	Pro	Asn	Lys	His	Gln	Ser	Ala
	450					455					460				
Val	Thr	Cys	Leu	Gln	Phe	Asn	Lys	Asn	Phe	Val	Ile	Thr	Ser	Ser	Asp
465					470					475					480
															-

Asp Gly Thr Val Lys Leu Trp Asp Leu Lys Thr Gly Glu Phe Ile Arg

485 490 495

Asn Leu Val Thr Leu Glu Ser Gly Gly Ser Gly Gly Val Val Trp Arg
500 505 510

Ile Arg Ala Ser Asn Thr Lys Leu Val Cys Ala Val Gly Ser Arg Asn
515 520 525

Gly Thr Glu Glu Thr Lys Leu Leu Val Leu Asp Phe Asp Val Asp Met 530 535 540

Lys

545

<210> 7

<211> 540

<212> PRT

<213> Homo sapiens

<400> 7

Met Lys Arg Lys Leu Asp His Gly Ser Glu Val Arg Ser Phe Ser Leu

1 5 10 15

Gly Lys Lys Pro Cys Lys Val Ser Glu Tyr Thr Ser Thr Thr Gly Leu
20 25 30

Val Pro Cys Ser Ala Thr Pro Thr Thr Phe Gly Asp Leu Arg Ala Ala 35 40 45

Asn Gly Gln Gln Gln Arg Arg Ile Thr Ser Val Gln Pro Pro 50 55 60

Thr Gly Leu Gln Glu Trp Leu Lys Met Phe Gln Ser Trp Ser Gly Pro 65 70 75 80

Glu Lys Leu	Leu Ala Leu	Asp Glu L	eu Ile Asp	Ser Cys Glu	Pro Thr
	85		90		95
Gln Val Lys	His Met Met	Gln Val I	le Glu Pro	Gln Phe Gln	Arg Asp
	100	1	.05	110	
Phe Ile Ser	Leu Leu Pro	Ive Clu L	eu Ala Leu	Tur Val Leu	Ser Dhe
	Deu Deu FIO	_	leu Ala Deu	_	Ser File
115		120		125	
Leu Glu Pro	Lys Asp Leu	Leu Gln A	la Ala Gln	Thr Cys Arg	Tyr Trp
130		135		140	
Arg Ile Leu	Ala Glu Asp	Asn Leu L	eu Trp Arg	Glu Lys Cys	Lys Glu
145	150		155		160
Glu Gly Ile	Asp Glu Pro	Leu His I	le Lys Arg	Arg Lys Val	Ile Lys
	165		170		175
Pro Gly Phe	Ile His Ser	Pro Trn I	ve Ser Ala	Tyr Ile Ara	Gln Hie
iio diy inc	180	_	185	190	0111 1125
	180	1	.63	150	
	_				_
Arg Ile Asp	Thr Asn Trp	Arg Arg G	Hy Glu Leu		Lys Val
195		200		205	
Leu Lys Gly	His Asp Asp	His Val I	Ile Thr Cys	Leu Gln Phe	Cys Gly
210		215		220	
Asn Arg Ile	Val Ser Gly	Ser Asp A	Asp Asn Thr	Leu Lys Val	Trp Ser
225	230	ı	235		240
Ala Val Thr	Gly Lys Cys	Leu Ara 1	Thr Leu Val	Glv His Thr	Glv Glv
/41 1111	245	1119	250	,	255
	44 3		250		233
			_	_	
Val Trp Ser	Ser Gln Met	Arg Asp A	Asn Ile Ile	Ile Ser Gly	Ser Thr
	260	2	265	270	

Asp Arg Thr Leu Lys Val Trp Asn Ala Glu Thr Gly Glu Cys Ile His

		275					280					285			
ጥኮ፦	Lou	Тиг	Glv	uic	Thr	Sar	ጥክν	Va 1	Ara	Cvc	Mot	Hic	Ī.au	Hic	

290 295 300

Lys Arg Val Val Ser Gly Ser Arg Asp Ala Thr Leu Arg Val Trp Asp 305 310 315 320

Ile Glu Thr Gly Gln Cys Leu His Val Leu Met Gly His Val Ala Ala 325 330 335

Val Arg Cys Val Gln Tyr Asp Gly Arg Arg Val Val Ser Gly Ala Tyr 340 345 350

Asp Phe Met Val Lys Val Trp Asp Pro Glu Thr Glu Thr Cys Leu His 355 360 365

Thr Leu Gln Gly His Thr Asn Arg Val Tyr Ser Leu Gln Phe Asp Gly 370 380

Ile His Val Val Ser Gly Ser Leu Asp Thr Ser Ile Arg Val Trp Asp 385 390 395 400

Val Glu Thr Gly Asn Cys Ile His Thr Leu Thr Gly His Gln Ser Leu 405 410 415

Thr Ser Gly Met Glu Leu Lys Asp Asn Ile Leu Val Ser Gly Asn Ala
420 425 430

Asp Ser Thr Val Lys Ile Trp Asp Ile Lys Thr Gly Gln Cys Leu Gln
435 440 445

Thr Leu Gln Gly Pro Asn Lys His Gln Ser Ala Val Thr Cys Leu Gln
450 455 460

Phe Asn Lys Asn Phe Val Ile Thr Ser Ser Asp Asp Gly Thr Val Lys

465 **4**70 **4**75 **4**80

Leu Trp Asp Leu Lys Thr Gly Glu Phe Ile Arg Asn Leu Val Thr Leu
485 490 495

Glu Ser Gly Gly Ser Gly Gly Val Val Trp Arg Ile Arg Ala Ser Asn 500 505 510

Thr Lys Leu Val Cys Ala Val Gly Ser Arg Asn Gly Thr Glu Glu Thr
515 520 525

Lys Leu Leu Val Leu Asp Phe Asp Val Asp Met Lys
530 535 540

<210> 8

<211> 589

<212> PRT

<213> Homo sapiens

<400> 8

Met Ser Lys Pro Gly Lys Pro Thr Leu Asn His Gly Leu Val Pro Val

1 5 10 15

Asp Leu Lys Ser Ala Lys Glu Pro Leu Pro His Gln Thr Val Met Lys
20 25 30

Ile Phe Ser Ile Ser Ile Ile Ala Gln Gly Leu Pro Phe Cys Arg Arg 35 40 45

Arg Met Lys Arg Lys Leu Asp His Gly Ser Glu Val Arg Ser Phe Ser 50 55 60

Leu Gly Lys Lys Pro Cys Lys Val Ser Glu Tyr Thr Ser Thr Thr Gly
65 70 75 80

Leu Val Pro Cys Ser Ala Thr Pro Thr Thr Phe Gly Asp Leu Arg Ala

				85					90					95	
Ala	Asn	Gly	Gln	Gly	Gln	Gln	Arg	Arg	Arg	Ile	Thr	Ser	Val	Gln	Pro
			100					105					110		
Dro	Шhr	Clv	Lou	Cln	Clu	Trn	Lou	Luc	Mot	Pho	Cln	Cor	m~n	Sor	Clu

Pro Thr Gly Leu Gln Glu Trp Leu Lys Met Phe Gln Ser Trp Ser Gly
115 120 125

Pro Glu Lys Leu Leu Ala Leu Asp Glu Leu Ile Asp Ser Cys Glu Pro 130 135 140

Asp Phe Ile Ser Leu Leu Pro Lys Glu Leu Ala Leu Tyr Val Leu Ser 165 170 175

Phe Leu Glu Pro Lys Asp Leu Leu Gln Ala Ala Gln Thr Cys Arg Tyr

180 185 190

Trp Arg Ile Leu Ala Glu Asp Asn Leu Leu Trp Arg Glu Lys Cys Lys
195 200 205

Glu Glu Gly Ile Asp Glu Pro Leu His Ile Lys Arg Arg Lys Val Ile 210 215 220

Lys Pro Gly Phe Ile His Ser Pro Trp Lys Ser Ala Tyr Ile Arg Gln 225 230 235 240

His Arg Ile Asp Thr Asn Trp Arg Arg Gly Glu Leu Lys Ser Pro Lys

245 250 255

Val Leu Lys Gly His Asp Asp His Val Ile Thr Cys Leu Gln Phe Cys
260 265 270

Gly Asn Arg Ile Val Ser Gly Ser Asp Asn Thr Leu Lys Val Trp

275 280 285

Thr Asp Arg Thr Leu Lys Val Trp Asn Ala Glu Thr Gly Glu Cys Ile
325 330 335

His Thr Leu Tyr Gly His Thr Ser Thr Val Arg Cys Met His Leu His 340 345 350

Glu Lys Arg Val Val Ser Gly Ser Arg Asp Ala Thr Leu Arg Val Trp
355 360 365

Asp Ile Glu Thr Gly Gln Cys Leu His Val Leu Met Gly His Val Ala 370 375 380

Ala Val Arg Cys Val Gln Tyr Asp Gly Arg Arg Val Val Ser Gly Ala 385 390 395 400

Tyr Asp Phe Met Val Lys Val Trp Asp Pro Glu Thr Glu Thr Cys Leu 405 410 415

His Thr Leu Gln Gly His Thr Asn Arg Val Tyr Ser Leu Gln Phe Asp 420 425 430

Gly Ile His Val Val Ser Gly Ser Leu Asp Thr Ser Ile Arg Val Trp
435 440 445

Asp Val Glu Thr Gly Asn Cys Ile His Thr Leu Thr Gly His Gln Ser
450 455 460

Leu Thr Ser Gly Met Glu Leu Lys Asp Asn Ile Leu Val Ser Gly Asn 465 470 475 480

Ala Asp Ser Thr Val Lys Ile Trp Asp Ile Lys Thr Gly Gln Cys Leu Gln Thr Leu Gln Gly Pro Asn Lys His Gln Ser Ala Val Thr Cys Leu Gln Phe Asn Lys Asn Phe Val Ile Thr Ser Ser Asp Asp Gly Thr Val Lys Leu Trp Asp Leu Lys Thr Gly Glu Phe Ile Arg Asn Leu Val Thr Leu Glu Ser Gly Gly Ser Gly Gly Val Val Trp Arg Ile Arg Ala Ser Asn Thr Lys Leu Val Cys Ala Val Gly Ser Arg Asn Gly Thr Glu Glu Thr Lys Leu Leu Val Leu Asp Phe Asp Val Asp Met Lys <210> 9 <211> 559 <212> PRT <213> Homo sapiens <400> 9 Met Lys Ile Phe Ser Ile Ser Ile Ile Ala Gln Gly Leu Pro Phe Cys

Phe Ser Leu Gly Lys Lys Pro Cys Lys Val Ser Glu Tyr Thr Ser Thr

Arg Arg Met Lys Arg Lys Leu Asp His Gly Ser Glu Val Arg Ser

35 40 45

Thr Gly Leu Val Pro Cys Ser Ala Thr Pro Thr Thr Phe Gly Asp Leu
50 55 60

Arg Ala Ala Asn Gly Gln Gly Gln Arg Arg Ile Thr Ser Val
65 70 75 80

Gln Pro Pro Thr Gly Leu Gln Glu Trp Leu Lys Met Phe Gln Ser Trp

85 90 95

Ser Gly Pro Glu Lys Leu Leu Ala Leu Asp Glu Leu Ile Asp Ser Cys
100 105 110

Glu Pro Thr Gln Val Lys His Met Met Gln Val Ile Glu Pro Gln Phe 115 · 120 125

Gln Arg Asp Phe Ile Ser Leu Leu Pro Lys Glu Leu Ala Leu Tyr Val 130 135 140

Leu Ser Phe Leu Glu Pro Lys Asp Leu Leu Gln Ala Ala Gln Thr Cys
145 150 155 160

Arg Tyr Trp Arg Ile Leu Ala Glu Asp Asn Leu Leu Trp Arg Glu Lys

165 170 175

Cys Lys Glu Glu Gly Ile Asp Glu Pro Leu His Ile Lys Arg Arg Lys

180 185 190

Val Ile Lys Pro Gly Phe Ile His Ser Pro Trp Lys Ser Ala Tyr Ile 195 200 205

Arg Gln His Arg Ile Asp Thr Asn Trp Arg Arg Gly Glu Leu Lys Ser 210 215 220

Pro Lys Val Leu Lys Gly His Asp Asp His Val Ile Thr Cys Leu Gln 225 230 235 240

	rue	cys	GTÅ	ASII	Arg	тте	vaı	ser	СТУ	ser	Asp	ASD	ASII	TILL	ьęи	пуѕ
					245					250					255	
,	Val	Trp	Ser	Ala	Val	Thr	Gly	Lys	Суѕ	Leu	Arg	Thr	Leu	Val	Gly	His
				260					265					270		
	Thr	Gly	Gly	Val	Trp	Ser	Ser	Gln	Met	Arg	Asp	Asn	Ile	Ile	Ile	Ser
			275					280					285			
	01	G	mla	2	3	mb	T	T	77a 1	M	2~~	21.	C1	mh	C1	C1
	GIY	Ser	rnr	Asp	Arg	'l'nr	Leu	гуѕ	Val	Trp	Asn	Ala	GIU	rnr	GIY	GIU
		290					295					300				
	Cve	Tla	ніс	Thr	T.eu	ጥህድ	Gly	His	Thr	Ser	Thr	Va1	Ara	Cvs	Met	His
		116	1115	1111	Бец		GLY	1115	1111	DCI		Vul	mrg	Cys	1100	
	305					310					315					320
	Leu	His	Glu	Lys	Arg	Val	Val	Ser	Gly	Ser	Arg	Asp	Ala	Thr	Leu	Arg
					225				_	330		_			225	
					325					330					335	
	Val	Trp	Asp	Ile	Glu	Thr	Gly	Gln	Cys	Leu	His	Val	Leu	Met	Gly	His
				340					345					350		
				340					343					330		
	Val	Ala	Ala	Val	Arg	Cys	Val	Gln	Tyr	Asp	Gly	Arg	Arg	Val	Val	Ser
			355					360					365			
	Gly	Ala	Tyr	Asp	Phe	Met	Val	Lys	Val	Trp	Asp	Pro	Glu	Thr	Glu	Thr
		370					375					380				
	_	_		 1	_	~ .	a 3		1	_	_		_	a	•	01 ···
	Cys	Leu	His	Thr	Leu	GIn	GIY	His	Thr	Asn	Arg	Val	'l'yr	Ser	Leu	Gln
	385					390					395					400
	Dh -	7	01	т1 ~	174 -	₹7- 1	₹7 ~ 1	0	01	0	T	7 ~~	መጐ ~	G~~	T1~	A ~~
	Lue	Asp	сту	тте	nis	vaı	vaı	ser	сту	ser	⊥eu	ASP	THE	ser	тте	Arg
					405					410					415	
	Va 1	ጥተኮ	Asn	Va 1	Glu	ጥ ኮ ጕ	Glv	Acn	Cve	Tle	Hic	Ψhr	Len	Thr	Glv	His
		11,0	1100		O.L.u		C-7	*****			****		204		- J	
				420					425					430		

Gly Asn Ala Asp Ser Thr Val Lys Ile Trp Asp Ile Lys Thr Gly Gln Cys Leu Gln Thr Leu Gln Gly Pro Asn Lys His Gln Ser Ala Val Thr Cys Leu Gln Phe Asn Lys Asn Phe Val Ile Thr Ser Ser Asp Asp Gly Thr Val Lys Leu Trp Asp Leu Lys Thr Gly Glu Phe Ile Arg Asn Leu Val Thr Leu Glu Ser Gly Gly Ser Gly Gly Val Val Trp Arg Ile Arg Ala Ser Asn Thr Lys Leu Val Cys Ala Val Gly Ser Arg Asn Gly Thr Glu Glu Thr Lys Leu Leu Val Leu Asp Phe Asp Val Asp Met Lys <210> 10 <211> 540 <212> PRT <213> Homo sapiens <400> 10 Met Lys Arg Lys Leu Asp His Gly Ser Glu Val Arg Ser Phe Ser Leu Gly Lys Lys Pro Cys Lys Val Ser Glu Tyr Thr Ser Thr Thr Gly Leu

Gln Ser Leu Thr Ser Gly Met Glu Leu Lys Asp Asn Ile Leu Val Ser

Val	Pro	Cys	Ser	Ala	Thr	Pro	Thr	Thr	Phe	Gly	Asp	Leu	Arg	Ala	Ala
		35					40					45			
Asn	Glv	Gln	Glv	Gln	Gln	Ara	Arq	Arq	Ile	Thr	Ser	Val	Gln	Pro	Pro
	50					55	J	,			60				
	30					,,,					•				
		_			_	_	_		-1	~3	_	_	~	01	_
	GLY	Leu	Gin	Glu	Trp	Leu	ьуs	Met	Pne		ser	Trp	ser	GIY	
65					70					75					80
Glu	Lys	Leu	Leu	Ala	Leu	Asp	Glu	Leu	Ile	Asp	Ser	Cys	Glu	Pro	Thr
				85					90					95	
Gln	Val	Lys	His	Met	Met	Gln	Val	Ile	Glu	Pro	Gln	Phe	Gln	Arg	Asp
			100					105					110		
Phe	Tle	Ser	Len	Len	Pro	Lvs	Glu	Len	Δla	ī _e n	Tvr	Val	Len	Ser	Phe
1110	110	115	Dea	Dea	110	275	120	200		204	-1-	125		551	10
		113					120					123			
							_								
Leu	Glu	Pro	Lys	Asp	Leu	Leu	Gln	Ala	Ala	Gln	Thr	Cys	Arg	Tyr	Trp
	130					135					140				
Arg	Ile	Leu	Ala	Glu	Asp	Asn	Leu	Leu	Trp	Arg	Glu	Lys	Cys	Lys	Glu
145					150					155					160
Glu	Gly	Ile	Asp	Glu	Pro	Leu	His	Ile	Lys	Arg	Arg	Lys	Val	Ile	Lys
				165					170					175	
Dro	C111	Dho	Tla	wi.c	Ser	Dro	m~n	Lvc	Sor	λ1 a	Път.	T1e	λrα	Gln	Hic
PIO	GIY	FIIE			ser	FIO	пр		Ser	Ala	ıyı	116		GIII	1115
			180					185					190		
Arg	Ile	Asp	Thr	Asn	Trp	Arg	Arg	Gly	Glu	Leu	Lys	Ser	Pro	Lys	Val
		195					200					205			
Leu	Lys	Gly	His	Asp	Asp	His	Val	Ile	Thr	Cys	Leu	Gln	Phe	Cys	Gly

Asn	Arg	Ile	Val	Ser	Gly	Ser	Asp	Asp	Asn	Thr	Leu	Lys	Val	Trp	Ser
225					230					235					240
Ala	Val	Thr	Gly	Lys	Cys	Leu	Arg	Thr	Leu	Val	Gly	His	Thr	Gly	Gly
				245					250					255	
Va1	Trp	Ser	Ser	Gln	Met	Arg	Asp	Asn	Ile	Ile	Ile	ser	Gly	Ser	Thr
			260					265					270		
Asp	Arg	Thr	Leu	Lys	Val	Trp	Asn	Ala	Glu	Thr	Gly	Glu	Cys	Ile	His
		275					280					285			
Thr	Leu	Tyr	Gly	His	Thr	Ser	Thr	Val	Arg	Cys	Met	His	Leu	His	Glu
	290					295					300				
Lys	Arg	Val	Val	Ser	Gly	Ser	Arg	Asp	Ala	Thr	Leu	Arg	Val	Trp	Asp
305					310					315					320
Ile	Glu	Thr	Gly	Gln	Cys	Leu	His	Val	Leu	Met	Gly	His	Val	Ala	Ala
			_	325	_				330		_			335	
Va1	Arg	Cys	Val	Gln	Tyr	Asp	Gly	Arg	Arg	Val	Val	Ser	Gly	Ala	Tyr
			340					345					350		
Asp	Phe	Met	Val	Lys	Val	Trp	Asp	Pro	Glu	Thr	Glu	Thr	Cys	Leu	His
		355					360					365			
Thr	Leu	Gln	Gly	His	Thr	Asn	Arg	Val	Tyr	Ser	Leu	Gln	Phe	Asp	Gly
	370					375			_		380				
Ile	His	Va1	Val	Ser	Gly	Ser	Leu	Asp	Thr	Ser	Ile	Arg	Val	Trp	Asp
385					390					395		J		-	400
Va1	Glu	Thr	Glv	Asn	Cvs	Ile	His	Thr	Leu	Thr	Glv	His	Gln	Ser	Leu
				405					410		2		-	415	-

Thr Ser Gly Met Glu Leu Lys Asp Asn Ile Leu Val Ser Gly Asn Ala
420 425 430

Asp Ser Thr Val Lys Ile Trp Asp Ile Lys Thr Gly Gln Cys Leu Gln
435 440 445

Thr Leu Gln Gly Pro Asn Lys His Gln Ser Ala Val Thr Cys Leu Gln
450 455 460

Phe Asn Lys Asn Phe Val Ile Thr Ser Ser Asp Asp Gly Thr Val Lys
465 470 475 480

Leu Trp Asp Leu Lys Thr Gly Glu Phe Ile Arg Asn Leu Val Thr Leu
485 490 495

Glu Ser Gly Gly Ser Gly Gly Val Val Trp Arg Ile Arg Ala Ser Asn 500 505 510

Thr Lys Leu Val Cys Ala Val Gly Ser Arg Asn Gly Thr Glu Glu Thr
515 520 525

Lys Leu Leu Val Leu Asp Phe Asp Val Asp Met Lys
530 535 540

<210> 11

<211> 34

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide primer

<400> 11

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<210>	12	
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<213>	Artificial Sequence	
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	Oligonucleotide primer	
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	Oligonucleotide primer	
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ggtaa	ttaca agttcttgtt gaactg	26

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	Artificial Sequence	
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	Oligonucleotide primer	
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<211> 1881
<212> DNA
<213> Artificial Sequence
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<223> Description of Artificial Sequence: 6 myc tagged
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gagcaaaagc tcatttctga agaggacttg aatgaaatgg agcaaaagct catttctgaa 180

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qqtqqagtat ggtcatcaca aatgagggac aacatcatca ttagtggatc tacagatcgg 1080
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tccatccgtg tttgggatgt ggagacaggg aattgcattc acacgttaac agggcaccag 1500
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aagcatcaga gtgctgtgac ctgtttacag ttcaacaaga actttgtaat taccagctca 1680
gatgatggaa ctgtaaaact atgggacttg aaaacgggtg aatttattcg aaacctagtc 1740
acattggaga gtgggggag tgggggagtt gtgtggcgga tcagagcctc aaacacaaag 1800
ctggtgtgtg cagttgggag tcggaatggg actgaagaaa ccaagctgct ggtgctggac 1860
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<210> 21
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<220>

<211> 626

<212> PRT

<213> Artificial Sequence

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Lvs	T.eu	Tle	Ser	G111	Glu	Asn	T. e 11	Asn	Glu	Met	Glu	Gln	LVS	T.e11	Tle
د رد	LCu	-1-0		Olu	014	1100	Lou		014	1100	014	0111		100	110
			20					25					30		
Ser	Glu	Glu	Asp	Leu	Asn	Glu	Met	Glu	Gln	Lys	Leu	Ile	Ser	Glu	Glu
		35					40					45			
Asp	Leu	Asn	Glu	Met	Glu	Gln	Lys	Leu	Ile	Ser	Glu	Glu	Asp	Leu	Asn
	50					55					60				
C111	Mot	C1.,	Cox	Tou	C111	7.00	Ton	mb ~	Mot	C111	C1n	Twa	T 011	T10	Cor
	Mec	GIU	ser	ьeu	Gly	ASP	ьeu	THE	Mec		GIN	гуѕ	ьеи	ire	
65					70					75					80
Glu	Glu	Asp	Leu	Asn	Ser	Met	Lys	Arg	Lys	Leu	Asp	His	Gly	Ser	Glu
				85					90					95	
Val	Arg	Ser	Phe	Ser	Leu	Gly	Lys	Lys	Pro	Cys	Lys	Val	Ser	Glu	Tyr
			100					105					110		
mb	Com	mla sa	mb	C1	T	77-7	Duc	C	0	21.	mh	Dwa	mb	mb	Dh a
THE	Ser		THE	GIY	ьeu	vaı		_	ser	АТА	THE		THE	THE	Phe
		115					120					125			
Gly	Asp	Leu	Arg	Ala	Ala	Asn	Gly	Gln	Gly	Gln	Gln	Arg	Arg	Arg	Ile
	130					135					140				
Thr	Ser	Val	Gln	Pro	Pro	Thr	Gly	Leu	Gln	Glu	Trp	Leu	Lys	Met	Phe
145					150					155	•		-		160
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<223> Description of Artificial Sequence: 6 myc tagged

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110	0.2.1.	195	022	9			200					205			
		132					200					205			
Leu	Tyr	Val	Leu	Ser	Phe	Leu	Glu	Pro	Lys	Asp	Leu	Leu	Gln	Ala	Ala
	210					215					220				
Gln	Thr	Cys	Arg	Tyr	Trp	Arg	Ile	Leu	Ala	Glu	Asp	Asn	Leu	Leu	Trp
225					230					235					240
Δτα	Glu	T.ve	Cys	Lvc	Glu	Glu	Glv	Tle	Asn	Glu	Pro	T.e.11	His	Tle	LVS
nrg	Giu	цys	СУЗ	Ī	GIU	Giu	Gly	110	_	O-u	110	пса	1115		цуз
				245					250					255	
Arg	Arg	Lys	Val	Ile	Lys	Pro	Gly	Phe	Ile	His	Ser	Pro	Trp	Lys	Ser
			260					265					270		
Ala	Tyr	Ile	Arg	Gln	His	Arg	Ile	Asp	Thr	Asn	Trp	Arg	Arg	Gly	Glu
		275					280					285			
T.eu	LVS	Ser	Pro	Lvs	Va l	Leu	LVS	Glv	His	Asp	Asp	His	Val	Tle	Thr
	290	501		-,,		295	_10	0-1			300				
	250					275					300				
Cys	Leu	Gln	Phe	Cys	Gly	Asn	Arg	Ile	Val	Ser	Gly	Ser	Asp	Asp	Asn
305					310					315					320
Thr	Leu	Lys	Val	Trp	Ser	Ala	Val	Thr	Gly	Lys	Cys	Leu	Arg	Thr	Leu
				325					330					335	
Val	Glv	His	Thr	Glv	Glv	Val	Trp	Ser	Ser	Gln	Met	Ara	qzA	Asn	Ile
	-2	_~	340		-2		-12	345		-3-		J	350		
			240					243					5 50		
													_		
Ile	Ile	Ser	Gly	Ser	Thr	Asp	Arg	Thr	Leu	Lys	Val	Trp	Asn	Ala	Glu
		355	,				360					365			

Thr	Gly	Glu	Cys	Ile	His	Thr	Leu	Tyr	Gly	His	Thr	Ser	Thr	Val	Arg
	370					375					380				
Суѕ	Met	His	Leu	His	Glu	Lys	Arg	Val	Val	Ser	Gly	Ser	Arg	Asp	Ala
385					390					395					400
Thr	Leu	Arg	Val	Trp	Asp	Ile	Glu	Thr	Gly	Gln	Cys	Leu	His	Val	Leu
				405					410					415	
Met	Gly	His	Val	Ala	Ala	Val	Arg	Cys	Val	Gln	Tyr	Asp	Gly	Arg	Arg
	-		420					425					430		
Val	Val	Ser	Glv	Ala	Tvr	qzA	Phe	Met	Val	Lvs	Val	Trp	Asp	Pro	Glu
		435	•		-	-	440			•		445	_		
Thr	Glu	Thr	Cvs	Leu	His	Thr	Leu	Gln	Glv	His	Thr	Asn	Arg	Val	Tyr
	450		_			455			_		460		J		-
Ser	Leu	Gln	Phe	Asp	Glv	Ile	His	Val	Val	Ser	Glv	Ser	Leu	Asp	Thr
465				•	470					475	•			-	480
Ser	Ile	Ara	Val	Trp	asp	Val	Glu	Thr	Glv	Asn	Cvs	Ile	His	Thr	Leu
		3		485					490					495	
Thr	Glv	His	Gln	Ser	Leu	Thr	Ser	Glv	Met	Glu	Leu	Lvs	Asp	Asn	Ile
			500					505					510		
Leu	Val	Ser	Glv	Asn	Ala	Asp	Ser	Thr	Val	Lvs	Ile	Trp	Asp	Ile	Lys
		515	-			-	520			•		525	•		-
							_ ^								
Thr	Gly	Gln	Cvs	Leu	Gln	Thr	Leu	Gln	Glv	Pro	Asn	Lvs	His	Gln	Ser
_ 	530		-1-2			535			1		540	_1 _			
	-55										- 10				
Ala	Val	ሞኮ r	Cve	Leu	Gln	Phe	Asn	Lve	Asn	Phe	Va 1	Tle	Thr	Ser	Ser
545			~, 13		550	- 110	-1011	-,,5	-1011	555		-10		201	560
242					220					ررر					550

Asp Asp Gly Thr Val Lys Leu Trp Asp Leu Lys Thr Gly Glu Phe Ile

565 570 575

Arg Asn Leu Val Thr Leu Glu Ser Gly Gly Ser Gly Gly Val Val Trp
580 585 590

Arg Ile Arg Ala Ser Asn Thr Lys Leu Val Cys Ala Val Gly Ser Arg
595 600 605

Asn Gly Thr Glu Glu Thr Lys Leu Leu Val Leu Asp Phe Asp Val Asp 610 615 620

Met Lys

625

<210> 22

<211> 31

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide primer

<400> 22

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31

<210> 23

<211> 29

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide primer

ggaattcctt catgtccaca tcaaagtcc

29

<210> 24

<211> 2010

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: V5HIS tagged homo sapien

<400> 24

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<210> 25

<211> 669

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: V5HIS tagged
homo sapien

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Cys Leu Ser Met Ser Thr Leu Glu Ser Val Thr Tyr Leu Pro Glu Lys
35 40 45

Gly Leu Tyr Cys Gln Arg Leu Pro Ser Ser Arg Thr His Gly Gly Thr
50 55 60

Glu Ser Leu Lys Gly Lys Asn Thr Glu Asn Met Gly Phe Tyr Gly Thr
65 70 75 80

Leu	Lys	Met	Ile	Phe	Tyr	Lys	Met	Lys	Arg	Lys	Leu	Asp	His	Gly	Ser
				85					90					95	
Glu	Val	Arσ	Ser	Phe	Ser	Leu	Gly	Lvs	Lvs	Pro	Cvs	Lvs	Va1	Ser	Glu
Oru	var	9			501		CII		בינם	120	010	2,0		50.	
			100					105					110		
Tyr	Thr	Ser	Thr	Thr	Gly	Leu	Val	Pro	Cys	Ser	Ala	Thr	Pro	Thr	Thr
		115					120					125			
Phe	Glv	Asp	Leu	Arq	Ala	Ala	Asn	Gly	Gln	Gly	Gln	Gln	Arq	Arq	Arq
	130	-		_		135		_		_	140		_		
	130					133					140				
Ile	Thr	Ser	Val	Gln	Pro	Pro	Thr	Gly	Leu	Gln	Glu	Trp	Leu	Lys	Met
145					150					155					160
Phe	Gln	Ser	Trp	Ser	Gly	Pro	Glu	Lys	Leu	Leu	Ala	Leu	Asp	Glu	Leu
				165					170					175	
Ile	Asp	Ser	Cys	Glu	Pro	Thr	Gln	Val	Lys	His	Met	Met	Gln	Val	Ile
			180					185					190		
Glu	Pro	Gln	Phe	Gln	Arg	Asp	Phe	Ile	Ser	Leu	Leu	Pro	Lys	Glu	Leu
		195					200					205			
	_	_		_	_	_,	_	~1	_	_	_	_	_	~1	
Ala	Leu	Tyr	Val	Leu	Ser	Pne	Leu	Glu	Pro	ьўs	Asp	Leu	Leu	GIn	Ala
	210					215					220				
Ala	Gln	Thr	Cys	Arg	Tyr	Trp	Arg	Ile	Leu	Ala	Glu	Asp	Asn	Leu	Leu
225					230					235					240
П	λ	C1	T	C	T	C11-	C1	C1	T1.	λ ~~	C1	Dws	T av-	u:-	T1 ^
Trb	AIG	GIU	ьуѕ		гуѕ	GIU	Glu	GIÀ		ASD	GIU	PLO	ьец		TIG
				245					250					255	
Lys	Arg	Arg	Lys	Val	Ile	Lys	Pro	Gly	Phe	Ile	His	Ser	Pro	Trp	Lys
			260					265					270		

Ser Ala Tyr Ile Arg Gln His Arg Ile Asp Thr Asn Trp Arg Arg Gly

275	280	285

Glu	Leu	Lys	Ser	Pro	Lys	Val	Leu	Lys	Gly	His	Asp	Asp	His	Val	Ile
	290					295					300				
Thr	Cys	Leu	Gln	Phe	Cys	Gly	Asn	Arg	Ile	Val	Ser	Gly	Ser	Asp	Asp
305					310					315					320
Asn	Thr	Leu	Lys	Val	Trp	Ser	Ala	Val	Thr	Gly	Lys	Cys	Leu	Arg	Thr
				325					330					335	
Leu	Val	Gly	His	Thr	Gly	Gly	Val	Trp	Ser	Ser	Gln	Met	Arg	Asp	Asn
		_	340		_			345					350		
Ile	Ile	Ile	Ser	Glv	Ser	Thr	Asp	Ara	Thr	Leu	Lvs	Val	Trp	Asn	Ala
		355		•			360	,				365	-		
Glu	ጥhr	Glv	Glu	Cvs	Tle	His	ጥhr	Len	Tur	Glv	His	Thr	Ser	Thr	Val
	370	0-1	J	0,0		375			-1-	J_1	380				
	,					3,3					300				
λrα	Cvc	Mot	Hic	LAU	шic	Glu	Lvc	Ara	Wal.	Val	Sor	Glu	Ser	Ara	Acn
385	СуБ	Mec	1115	Dea	390	GIU	пуъ	ALG	Val	395	Ser	GLY	261	ALG	400
303					390					333					400
7 .1	mb	T	3	₹ 7 - 1	П	7	т1 -	01 11	mb	01	01n	O	7	uic	17-1
Ala	Thr	Leu	Arg		Trp	Asp	iie	GIU		GIY	GIn	Cys	Leu		vaı
				405					410					415	
_				1			•	_				_	_		
Leu	Met	GLy		Val	Ala	Ala	Val		Cys	Val	Gln	Tyr	Asp	GŢĀ	Arg
			420					425					430		

Glu Thr Glu Thr Cys Leu His Thr Leu Gln Gly His Thr Asn Arg Val
450 455 460

Arg Val Val Ser Gly Ala Tyr Asp Phe Met Val Lys Val Trp Asp Pro

445

440

435

Tyr Ser Leu Gln Phe Asp Gly Ile His Val Val Ser Gly Ser Leu Asp

Thr	Ser	Ile	Arg	Val 485	Trp	Asp	Val	Glu	Thr 490	Gly	Asn	Cys	Ile	His 495	Thr
Leu	Thr	Gly	His 500	Gln	Ser	Leu	Thr	Ser 505	Gly	Met	Glu	Leu	Lys 510	Asp	Asn
Ile	Leu	Val 515	Ser	Gly	Asn	Ala	Asp 520	Ser	Thr	Val	Lys	Ile 525	Trp	Asp	Ile
Lys	Thr 530	Gly	Gln	Cys	Leu	Gln 535	Thr	Leu	Gln	Gly	Pro 540	Asn	Lys	His	Gln
Ser 545	Ala	Val	Thr	Cys	Leu 550	Gln	Phe	Asn	Lys	Asn 555	Phe	Val	Ile	Thr	Ser 560
Ser	Asp	Asp	Gly	Thr 565	Val	Lys	Leu	Trp	Asp 570	Leu	Lys	Thr	Gly	Glu 575	Phe
Ile	Arg	Asn	Leu 580	Val	Thr	Leu	Glu	Ser 585	Gly	Gly	Ser	Gly	Gly 590	Val	Val
Trp	Arg	Ile 595	Arg	Ala	Ser	Asn	Thr 600	Lys	Leu	Val	Cys	Ala 605	Val	Gly	Ser
Arg	Asn 610	Gly	Thr	Glu	Glu	Thr 615	Lys	Leu	Leu	Val	Leu 620	Asp	Phe	Asp	Val
625		_			630		_			635					Glu 640
Ser	Arg	Gly	Pro	Phe		Gly	Lys	Pro	Ile 650	Pro	Asn	Pro	Leu	Leu 655	Gly

Leu Asp Ser Thr Arg Thr Gly His His His His His

<210> 26

<211> 2001

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: MYCHIS tagged
homo sapiens

<400> 26

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<210> 27

<211> 666

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: MYCHIS tagged homo sapiens

<400> 27

Met Cys Val Pro Arg Ser Gly Leu Ile Leu Ser Cys Ile Cys Leu Tyr

1 5 10 15

Cys Gly Val Leu Leu Pro Val Leu Leu Pro Asn Leu Pro Phe Leu Thr

Cys Leu Ser Met Ser Thr Leu Glu Ser Val Thr Tyr Leu Pro Glu Lys 35 40 45

Gly Leu Tyr Cys Gln Arg Leu Pro Ser Ser Arg Thr His Gly Gly Thr
50 55 60

Glu Ser Leu Lys Gly Lys Asn Thr Glu Asn Met Gly Phe Tyr Gly Thr
65 70 75 80

Leu Lys Met Ile Phe Tyr Lys Met Lys Arg Lys Leu Asp His Gly Ser

85	90	95

Glu	Val	Arg	Ser	Phe	Ser	Leu	Gly	Lys	Lys	Pro	Cys	Lys	Val	Ser	Glu
			100					105					110		
Tyr	Thr	Ser	Thr	Thr	Gly	Leu	Val	Pro	Cys	Ser	Ala	Thr	Pro	Thr	Thr
		115					120					125			
Phe	Gly	Asp	Leu	Arg	Ala	Ala	Asn	Gly	Gln	Gly	Gln	Gln	Arg	Arg	Arg
	130					135					140				

Ile Thr Ser Val Gln Pro Pro Thr Gly Leu Gln Glu Trp Leu Lys Met
145 150 155 160

Phe Gln Ser Trp Ser Gly Pro Glu Lys Leu Leu Ala Leu Asp Glu Leu
165 170 175

Ile Asp Ser Cys Glu Pro Thr Gln Val Lys His Met Met Gln Val Ile 180 185 190

Glu Pro Gln Phe Gln Arg Asp Phe Ile Ser Leu Leu Pro Lys Glu Leu
195 200 205

Ala Leu Tyr Val Leu Ser Phe Leu Glu Pro Lys Asp Leu Leu Gln Ala 210 215 220

Ala Gln Thr Cys Arg Tyr Trp Arg Ile Leu Ala Glu Asp Asn Leu Leu 225 230 235 240

Trp Arg Glu Lys Cys Lys Glu Glu Gly Ile Asp Glu Pro Leu His Ile
245 250 255

Lys Arg Arg Lys Val Ile Lys Pro Gly Phe Ile His Ser Pro Trp Lys
260 265 270

Ser Ala Tyr Ile Arg Gln His Arg Ile Asp Thr Asn Trp Arg Arg Gly

Glu	Leu	Lys	Ser	Pro	Lys	Val	Leu	Lys	Gly	His	Asp	Asp	His	Val	Ile
	290					295					300				
Thr	Cvs	Leu	Gln	Phe	Cvs	Glv	Asn	Ara	Ile	Val	Ser	Glv	Ser	Asp	Asp
305	-1-				310		-	J		315		-	_	-	320
303					310					313					320
	m)	_	_	**- 7		a		**. 1	m1	01	•	~	•	•	ml
Asn	unr	Leu	Lys		Trp	ser	Ala	vai		GIY	ьуs	Cys	ьeu		Thr
				325					330					335	
Leu	Val	Gly	His	Thr	Gly	Gly	Val	Trp	Ser	Ser	Gln	Met	Arg	Asp	Asn
			340					345					350		
Ile	Ile	Ile	Ser	Gly	Ser	Thr	Asp	Arg	Thr	Leu	Lys	Val	Trp	Asn	Ala
		355					360					365			
Glu	Thr	Gly	Glu	Cys	Ile	His	Thr	Leu	Tyr	Gly	His	Thr	Ser	Thr	Val
	370					375					380				
λτα	Cvc	Mot	His	T.011	ніс	Glu	Luc	Δνα	Va l	Ual	Sar	Gly	Sar	Δνα	λen
385	Суз	nec	1113	Бец	390	GIU	цуз	Arg	vai	395	Jei	GLY	Jei	nrg	
303					390					393					400
								_							
Ala	Thr	Leu	Arg	Val	Trp	Asp	Ile	Glu		Gly	Gln	Cys	Leu	His	Val
				405					410					415	
Leu	Met	Gly	His	Val	Ala	Ala	Val	Arg	Cys	Val	Gln	Tyr	Asp	Gly	Arg
			420					425					430		
Arg	Val	Val	Ser	Gly	Ala	Tyr	Asp	Phe	Met	Val	Lys	Val	Trp	Asp	Pro
		435					440					445			
Glu	Thr	Glu	Thr	Cys	Leu	His	Thr	Leu	Gln	Gly	His	Thr	Asn	Arg	Val
	450					455				-	460				
Паг⊶	50×	Lou	Gln	Dho	λ o.~	C1	т1-	ui.	17a 1	77-7	Cor.	C1.	505	Lou	∆ c~
тĀт	Ser	neu	3111	1-11G	ηsp	GT A	T T G	1112	val	val	Set	GTA	SET	пeп	uah

Thr	Ser	Ile	Arg	Val 485	Trp	Asp	Val	Glu	Thr 490	Gly	Asn	Cys	Ile	His 495	Thr
Leu	Thr	Gly	His 500	Gln	Ser	Leu	Thr	Ser 505	Gly	Met	Glu	Leu	Lys 510	Asp	Asn
Ile	Leu	Val 515	Ser	Gly	Asn	Ala	Asp 520	Ser	Thr	Val	Lys	Ile 525	Trp	Asp	Ile
Lys	Thr 530	Gly	Gln	Cys	Leu	Gln 535	Thr	Leu	Gln	Gly	Pro 540	Asn	Lys	His	Gln
Ser 545	Ala	Val	Thr	Суз	Leu 550	Gln	Phe	Asn	Lys	Asn 555	Phe	Val	Ile	Thr	Ser 560
Ser	Asp	Asp	Gly	Thr 565	Val	Lys	Leu	Trp	Asp 570	Leu	Lys	Thr	Gly	Glu 575	Phe
Ile	Arg	Asn	Leu 580	Val	Thr	Leu	Glu	Ser 585	Gly	Gly	Ser	Gly	Gly 590	Val	Val
Trp	Arg	Ile 595	Arg	Ala	Ser	Asn	Thr	Lys	Leu	Val	Cys	Ala 605	Val	Gly	Ser
Arg	Asn 610	Gly	Thr	Glu	Glu	Thr	Lys	Leu	Leu	Val	Leu 620	Asp	Phe	Asp	Val
Asp 625	Met	Lys	Glu	Phe	Cys 630	Arg	Tyr	Pro	Ala	Gln 635	Trp	Arg	Pro	Leu	Glu 640
Ser	Arg	Gly	Pro	Phe	Glu	Gln	Lys	Leu	Ile 650	Ser	Glu	Glu	Asp	Leu 655	Asn
Met	His	Thr	Gly	His	His	His	His	His	His						